

Table S5. TargetScanS predicts that miR-34a interacts with *Atg9A* 3'-UTR

Gene	Position	Predicted consequential pairing of target region (top) and miRNA (bottom)	Seed match
<i>ATG9A</i>	285-291 of 3'-UTR	5' ...CAGCCCAGCCCCAGUACUGGCCAC... 	7mer-1A
<i>hsa-miR-34a</i>		3' UGUUGGGUCGAUUCUGUGACGGU	